

SEQUENCE LISTING

<110> Burnie, James  
Matthews, Ruth C.

<120> Treatment of Infection Due to Clostridium Difficile

<130> 3022-109

<140> 10/550,410  
<141> 2005-09-23

<150> PCT/GB2004/001383  
<151> 2004-03-25

<150> GB0306782.4  
<151> 2003-03-25

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<170> PatentIn version 3.3

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Phe Gly Ala Arg Asp Tyr Glu Glu Pro Val Ile Lys Lys Trp Ser Glu	
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gaa cat aag gat gtt caa gtg gat att tat cct gaa aac atg act gaa	150
Glu His Lys Asp Val Gln Val Asp Ile Tyr Pro Glu Asn Met Thr Glu	
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gaa aat gta gtt aaa gct aaa ggg tat gat ggt ata tct ata caa caa	198
Glu Asn Val Val Lys Ala Lys Gly Tyr Asp Gly Ile Ser Ile Gln Gln	
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act aac tat ata gat aat cct tat att tat gaa act tta aaa gat gct	246
Thr Asn Tyr Ile Asp Asn Pro Tyr Ile Tyr Glu Thr Leu Lys Asp Ala	
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ggg gtt aaa gtt ata gct tca aga act gca ggg gtt gac atg ata cat	294
Gly Val Lys Val Ile Ala Ser Arg Thr Ala Gly Val Asp Met Ile His	

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tat tca cct aat gca ata gct gaa tta gct gtt act caa gct atg aac Tyr Ser Pro Asn Ala Ile Ala Glu Leu Ala Val Thr Gln Ala Met Asn 105	110	115		390
ctt tta aga aag act cct cta gta aag aaa aaa gtc tgt gaa ggt gat Leu Leu Arg Lys Thr Pro Leu Val Lys Lys Lys Val Cys Glu Gly Asp 120	125	130		438
tac cgt tgg ata gct gaa ctt ctt gga aca gaa gtt aga tct att aca Tyr Arg Trp Ile Ala Glu Leu Leu Gly Thr Glu Val Arg Ser Ile Thr 135	140	145		486
gtt ggt gtt ata ggt aca gga aaa ata ggt gct act tct gca aaa tta Val Gly Val Ile Gly Thr Gly Lys Ile Gly Ala Thr Ser Ala Lys Leu 150	155	160	165	534
ttc aaa ggc cta gga gct aat gta att gca tat gac caa tat cca aat Phe Lys Gly Leu Gly Ala Asn Val Ile Ala Tyr Asp Gln Tyr Pro Asn 170	175	180		582
agt gat tta aac gat ata tta act tac aaa gat tct tta gaa gac ctt Ser Asp Leu Asn Asp Ile Leu Thr Tyr Lys Asp Ser Leu Glu Asp Leu 185	190	195		630
cta aaa gaa gct gac gtt ata aca tta cat act cct tta ctt gaa gga Leu Lys Glu Ala Asp Val Ile Thr Leu His Thr Pro Leu Leu Glu Gly 200	205	210		678
aca aaa cat atg ata aat aaa gat act cta gct ata atg aag gat gga Thr Lys His Met Ile Asn Lys Asp Thr Leu Ala Ile Met Lys Asp Gly 215	220	225		726
gct tac ata gta aat act gcc cgt ggt ggt tta att gat aca ggg gat Ala Tyr Ile Val Asn Thr Ala Arg Gly Gly Leu Ile Asp Thr Gly Asp 230	235	240	245	774
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aca ttt gaa act gaa gga ttg ttc tta aac aaa aaa atg aat cct gga Thr Phe Glu Thr Glu Gly Leu Phe Leu Asn Lys Lys Met Asn Pro Gly 265	270	275		870
gaa tta act gac cca gaa ata aat aaa ctt ctt tct atg gaa caa gtt Glu Leu Thr Asp Pro Glu Ile Asn Lys Leu Leu Ser Met Glu Gln Val 280	285	290		918
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Ser Leu Glu Asp Leu Leu Lys Glu Ala Asp Val Ile Thr Leu His Thr  
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Pro Leu Leu Glu Gly Thr Lys His Met Ile Asn Lys Asp Thr Leu Ala  
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Ile Met Lys Asp Gly Ala Tyr Ile Val Asn Thr Ala Arg Gly Gly Leu  
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Ala Ala Ala Leu Asp Thr Phe Glu Thr Glu Gly Leu Phe Leu Asn Lys  
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